

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/903,823

DATE: 01/03/2002

TIME: 11:29:07

Input Set : N:\Crif3\RULE60\09903823.raw

Output Set: N:\CRF3\01032002\I903823.raw

RECEIVED

APR 01 2002

TECH CENTER 1600/2900

ENTERED

1 <110> APPLICANT: Genentech, Inc.
 2 Ashkenazi, Avi
 3 Botstein, David
 4 Desnoyers, Luc
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 6 Ferrara, Napoleone
 7 Filvaroff, Ellen
 8 Fong, Sherman
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 10 Gerber, Hanspeter
 11 Gerritsen, Mary E.
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 13 Godowski, Paul J.
 14 Grimaldi, Christopher J.
 15 Gurney, Austin L.
 16 Hillan, Kenneth, J.
 17 Kljavin, Ivar J.
 18 Mather, Jennie P.
 19 Pan, James
 20 Paoni, Nicholas F.
 21 Roy, Margaret Ann
 22 Stewart, Timothy A.
 23 Tumas, Daniel
 24 Williams, P. Mickey
 25 Wood, William, I.
 26 <120> TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
 27 Acids Encoding the Same
 28 <130> FILE REFERENCE: 10466-14
 29 <140> CURRENT APPLICATION NUMBER: 09/903,823
 30 <141> CURRENT FILING DATE: 2001-07-11
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 32 <151> PRIOR FILING DATE: 2000-09-18
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 34 <151> PRIOR FILING DATE: 1999-07-07
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 38 <151> PRIOR FILING DATE: 1999-07-28
 39 <150> PRIOR APPLICATION NUMBER: PCT/US99/20594
 40 <151> PRIOR FILING DATE: 1999-09-08
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 42 <151> PRIOR FILING DATE: 1999-09-13
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74      tggagctccg gctgcgtctt cccgcagcgc taccgcccat gcgcctgccg 150
75      cgccggggccg cgctggggct cctgccgctt ctgctgctgc tgccgcccgc 200
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78      ggcgggaaca cggcttgggg ggaaaagacg ctgtccaagt acgagtccag 350
79      cgagattcgc ctgctggaga tcctggaggg gctgtgagag agcagcgact 400
80      tcgaatgcaa tcagatgcta gaggcgcagg aggagcacct ggaggcctgg 450
81      tggctgcagc tgaagagcga atatcctgac ttattcgagt ggttttgtgt 500
82      gaagacactg aaagtgtgct gctctccagg aacctacggt cccgactgtc 550
83      tcgcatgcca gggcggatcc cagaggccct gcagcgggaa tggccactgc 600
84      agcggagatg ggagcagaca gggcgacggg tcctgccggt gccacatggg 650
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87      acgtgctcgg gcctgaccaa cagagactgc ggcgagtgtg aagtgggctg 800
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96      tgtgccggac ttacccttta aattattcag aaggatgtcc cgtggaaaat 1250
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100     aaaattgacc attgtaggta atcaggagga aaaaaaaaaa aaaaaaaaaa 1450
101     aaagggcggc cgcgactcta gagtcgacct gcagaagctt ggccgccatg 1500
102     gcccaacttg tttattgcag cttataatgg ttacaaataa agcaatagca 1550
103     tcacaaattt cacaaataaa gcattttttt cactgcattc tagttgtggt 1600
104     ttgtccaaac tcatcaatgt atcttatcat gtctggatcg ggaattaatt 1650
105     cggcgcagca ccatggcctg aaataacctc tgaaagagga acttggttag 1700
106     gtaccttctg aggcggaaag aaccagctgt ggaatgtgtg tcagttaggg 1750
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118             20             25             30
119     Cys His Arg Cys Arg Gly Leu Val Asp Lys Phe Asn Gln Gly Met
120             35             40             45
121     Val Asp Thr Ala Lys Lys Asn Phe Gly Gly Gly Asn Thr Ala Trp
122             50             55             60
123     Glu Glu Lys Thr Leu Ser Lys Tyr Glu Ser Ser Glu Ile Arg Leu
124             65             70             75
125     Leu Glu Ile Leu Glu Gly Leu Cys Glu Ser Ser Asp Phe Glu Cys
126             80             85             90
127     Asn Gln Met Leu Glu Ala Gln Glu Glu His Leu Glu Ala Trp Trp
128             95            100            105
129     Leu Gln Leu Lys Ser Glu Tyr Pro Asp Leu Phe Glu Trp Phe Cys
130            110            115            120
131     Val Lys Thr Leu Lys Val Cys Cys Ser Pro Gly Thr Tyr Gly Pro
132            125            130            135
133     Asp Cys Leu Ala Cys Gln Gly Gly Ser Gln Arg Pro Cys Ser Gly
134            140            145            150
135     Asn Gly His Cys Ser Gly Asp Gly Ser Arg Gln Gly Asp Gly Ser
136            155            160            165
137     Cys Arg Cys His Met Gly Tyr Gln Gly Pro Leu Cys Thr Asp Cys
138            170            175            180
139     Met Asp Gly Tyr Phe Ser Ser Leu Arg Asn Glu Thr His Ser Ile
140            185            190            195
141     Cys Thr Ala Cys Asp Glu Ser Cys Lys Thr Cys Ser Gly Leu Thr
142            200            205            210
143     Asn Arg Asp Cys Gly Glu Cys Glu Val Gly Trp Val Leu Asp Glu
144            215            220            225
145     Gly Ala Cys Val Asp Val Asp Glu Cys Ala Ala Glu Pro Pro Pro
146            230            235            240
147     Cys Ser Ala Ala Gln Phe Cys Lys Asn Ala Asn Gly Ser Tyr Thr

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150		260	265	270
151	Pro Gly Asn Cys	Lys Glu Cys Ile Ser	Gly Tyr Ala Arg Glu His	
152		275	280	285
153	Gly Gln Cys Ala	Asp Val Asp Glu Cys	Ser Leu Ala Glu Lys Thr	
154		290	295	300
155	Cys Val Arg Lys	Asn Glu Asn Cys Tyr	Asn Thr Pro Gly Ser Tyr	
156		305	310	315
157	Val Cys Val Cys	Pro Asp Gly Phe Glu	Glu Thr Glu Asp Ala Cys	
158		320	325	330
159	Val Pro Pro Ala	Glu Ala Glu Ala Thr	Glu Gly Glu Ser Pro Thr	
160		335	340	345
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166 <212> TYPE: DNA

167 <213> ORGANISM: Homo Sapien

168 <400> SEQUENCE: 3

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171   cgcccagccg tctaaacggg aacagccctg gctgagggag ctgcagcgca 150
172   gcagagtatc tgacggcgcc aggttgcgta ggtgcggcac gaggagtttt 200
173   cccggcagcg aggaggtcct gagcagcatg gcccgaggga gcgccttccc 250
174   tgccgcgcgc ctctggctct ggagcatcct cctgtgcctg ctggcactgc 300
175   gggcgagggc cgggcccgcg caggaggaga gcctgtacct atggatcgat 350
176   gctcaccagg caagagtact cataggattt gaagaagata tcctgattgt 400
177   ttcagagggg aaaatggcac cttttacaca tgatttcaga aaagcgcaac 450
178   agagaatgcc agctattcct gtcaatatcc attccatgaa ttttacctgg 500
179   caagctgcag ggcaggcaga atacttctat gaattcctgt ccttgcgctc 550
180   cctggataaa ggcacatcgg cagatccaac cgtcaatgtc cctctgctgg 600
181   gaacagtgcc tcacaaggca tcagttgttc aagttggttt cccatgtctt 650
182   ggaaaacagg atgggggtggc agcatttgaa gtggatgtga ttgttatgaa 700
183   ttctgaaggc aacaccattc tccaaacacc tcaaaatgct atcttcttta 750
184   aaacatgtca acaagctgag tgcccaggcg ggtgccgaaa tggaggcttt 800
185   tgtaatgaaa gacgcactct cgagtgtcct gatgggttcc acggacctca 850
186   ctgtgagaaa gccctttgta cccacgatg tatgaatggg ggactttgtg 900
187   tgactcctgg tttctgcac tgcccacctg gattctatgg agtgaactgt 950
188   gacaaagcaa actgctcaac cacctgcttt aatggaggga cctgtttcta 1000
189   ccctggaaaa tgtatttgcc ctccaggact agagggagag cagtgtgaaa 1050
190   tcagcaaata cccacaaccc tgtcgaaatg gaggtaaatg cattggtaaa 1100
191   agcaaatagt agtgttccaa aggttaccag ggagacctct gttcaaagcc 1150
192   tgtctgcgag cctggctgtg gtgcacatgg aacctgccat gaaccaaca 1200
193   aatgccaatg tcaagaaggt tggcatggaa gacactgcaa taaaagggtac 1250
194   gaagccagcc tcatacatgc cctgaggcca gcaggcgccc agctcaggca 1300
195   gcacacgcct tcacttaaaa aggccgagga gcggcgggat ccacctgaat 1350
196   ccaattacat ctggtgaact ccgacatctg aaacgtttta agttacacca 1400
197   agttcatagc ctttggttaac ctttcatgtg ttgaatgttc aaataatggt 1450

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Output Set: N:\CRF3\01032002\I903823.raw

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199   actgagctga tatttactct tccttttaag ttttctaagt acgtctgtag 1550
200   catgatggta tagattttct tgtttcagtg ctttgggaca gattttatat 1600
201   tatgtcaatt gatcagggtta aaattttcag tgtgtagttg gcagatatatt 1650
202   tcaaaattac aatgcattta tgggtgtctgg gggcagggga acatcagaaa 1700
203   ggttaaattg ggcaaaaatg cgtaagtcac aagaatttgg atggtgcagt 1750
204   taatgttgaa gttacagcat ttcagatttt attgtcagat atttagatgt 1800
205   ttgttacatt tttaaaaatt gctcttaatt tttaaactct caatacaata 1850
206   tattttgacc ttaccattat tccagagatt cagtattaaa aaaaaaaaaa 1900
207   ttacactgtg gtagtggcat ttaaacaata taatatattc taaacacaat 1950
208   gaaataggga atataatgta tgaacttttt gcattggctt gaagcaatat 2000
209   aatatattgt aaacaaaaca cagctcttac ctaataaaca ttttatactg 2050
210   tttgtatgta taaaataaag gtgctgcttt agtttttttg aaaaaaaaaa 2100
211   aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa gggcggccgc gactctagag 2150
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223               20               25               30
224   Pro Gln Glu Glu Ser Leu Tyr Leu Trp Ile Asp Ala His Gln Ala
225               35               40               45
226   Arg Val Leu Ile Gly Phe Glu Glu Asp Ile Leu Ile Val Ser Glu
227               50               55               60
228   Gly Lys Met Ala Pro Phe Thr His Asp Phe Arg Lys Ala Gln Gln
229               65               70               75
230   Arg Met Pro Ala Ile Pro Val Asn Ile His Ser Met Asn Phe Thr
231               80               85               90
232   Trp Gln Ala Ala Gly Gln Ala Glu Tyr Phe Tyr Glu Phe Leu Ser
233               95              100              105
234   Leu Arg Ser Leu Asp Lys Gly Ile Met Ala Asp Pro Thr Val Asn
235               110             115             120
236   Val Pro Leu Leu Gly Thr Val Pro His Lys Ala Ser Val Val Gln
237               125             130             135
238   Val Gly Phe Pro Cys Leu Gly Lys Gln Asp Gly Val Ala Ala Phe
239               140             145             150
240   Glu Val Asp Val Ile Val Met Asn Ser Glu Gly Asn Thr Ile Leu
241               155             160             165
242   Gln Thr Pro Gln Asn Ala Ile Phe Phe Lys Thr Cys Gln Gln Ala
243               170             175             180
244   Glu Cys Pro Gly Gly Cys Arg Asn Gly Gly Phe Cys Asn Glu Arg
245               185             190             195
246   Arg Ile Cys Glu Cys Pro Asp Gly Phe His Gly Pro His Cys Glu
247               200             205             210

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/903,823

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